

## SEQUENCE LISTING PART OF THE DESCRIPTION

## pONY8.1Z MLVHyb (SEQ ID NO 10)

5 AGATCTTGAATAATAAAATGTGTGTTTGTCCGAAATACGCGTTTTGAGATTTCTGTCGCCGACTAAATTCATGTCCGCCG  
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### pONY8.3G FB29 – (SEQ ID No 45)

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**pONY8.3G FB29 + (SEQ ID No 46)**

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**pONY8.3GPGK – (SEQ ID No 47)**

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**pONY8.3G PGK + (SEQ ID No 48)**

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## pONY3.2IREShyg

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**pONY8ZA CMVHyb (SEQ ID N 52)**

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**PEsynGP (SEQ ID No 53)**

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5 GTCCGGTGCCCTGAATGAACTGCAGGACGAGGCAGCGCGGCTATCGTGGCTGGCCACGAC  
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**PESDSYNGP (SEQ ID No 54)**

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**MLV construct CZCG (SEQ ID No 55)**

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GGTGCCCCCAAGGACCTGAAATGACCCTGTGCCTTATTTGAACTAACCAATCAGTTCGCTT  
CTCGCTTCTGTTTCGCGCGCTTCTGCTCCCCGAGCTCAATAAAAGAGCCCAACCCCTCA  
CTCGGGGCGCCAGTCCCTCCGATTGACTGAGTCGCCCCGGGTACCCGTGTATCCAATAAACC  
35 CTCTTGCAAGTGCATCCGACTTGTGGTCTCGCTGTTTCTTGGGAGGGTCTCCTCTGAGTG  
ATTGACTACCCGTGACGGGGGTCTTTTCAATTTGGGGGCTCGTCCGGGATCGGGAGACCCC  
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CGGATGCCGGGAGCAGACAAGCCGTCAGGGCGCGTCAGCGGGTGTGGCGGGTGTGCGG  
40 GCGCAGCCATGACCCAGTCACGTAGCGATAGCGGAGTGATACTGGCTTAAGTATGCGGC  
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GGTCTGTTGCGGTGCGGCGAGCGGTATCAGCTCACTCAAAGGCGGTAATACGGTTATCCAC  
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45 CCGTAAAAAGGCCGCGTTTGCTGGCGTTTTTCCATAGGCTCCGCCCCCTGACGAGCATCA  
CAAAAAATCGACGCTCAAGTCAGAGGTGGCGAAACCCGACAGGACTATAAAGATACCAGGC  
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50 GCCCGACCGCTGCGCCTTATCCGGTAACTATCGTCTTGAGTCCAACCCGGTAAGACACGA  
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CAAACAAACCACCGCTGGTAGCGGTGGTTTTTTTGTGTTGCAAGCAGCAGATTACGCGCAG  
55 AAAAAAGGATCTCAAGAAGATCCTTTGATCTTTTCTACGGGTCTGACGCTCAGTGGAA  
CGAAACTCACGTTAAGGGATTTTGGTCATGAGATTATCAAAAAGGATCTTACCTAGAT  
CCTTTTAAATTAATAATGAAGTTTAAATCAATCTAAAGTATATATGAGTAACTTGGTC  
TGACAGTTACCAATGCTTAATCAGTGAGGCACCTATCTCAGCGATCTGTCTATTTTCGTTT  
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60 TGGCCCCAGTGCTGCAATGATACCGCGAGACCCACGCTCACCGGCTCCAGATTTATCAGC

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5 AAAAGCGGTTAGCTCCTTCGGTCTCCGATCGTTGTCAGAAGTAAGTTGGCCGAGTGT  
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10 GAGATCCAGTTCGATGTAACCCACTCGTGCACCCAACTGATCTTCAGCATCTTTACTTT  
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15 CATGACATTAACCTATAAAAATAGGCGTATCACGAGGCCCTTTCGTCTCAAGAATTCAT  
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20 TCGTCCAACCAACCGACTCTGACGGCAGTTTACGAGAGAGATGATAGGGTCTGCTCAG  
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25 TCCCATAGTAACGCCAATAGGGACTTTCCATTGACGTCAATGGGTGGAGTATTTACGGTA  
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30 TGACGTCAATGGGAGTTTGTTTTGGCACCAAAATCAACGGGACTTTCCAAAATGTCGTAA  
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35 GATTTGGAGACCCCTGCCAGGGACCACCGACCCACCACCGGGAGGTAAGCTGGCCAGCA  
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40 GACGAGAACCATAAACAGTTCCCGCCTCCGTCTGAATTTTGTCTTTCGGTTTGGAAACCGA  
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GTGTTTCTGTATTTGTCTGAAAATTAGGGCCAGACTGTTACCACTCCCTTAAGTTTGACC  
TTAGGTCACTGGAAGATGTGAGCGGATCGCTCACAACCAGTCGGTAGATGTCAAGAAG  
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45

**PCGCLNGFR (SEQ ID No 57)**

GTTACCTTCTGCTCTGCAGAATGGCCAACCTTTAACGTCCGATGGCCGCGAGACGGCACC  
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50 CACCCAGACCAGGTCCCTACATCGTGACCTGGGAAGCCTTGGCTTTTGACCCCCCTCCC  
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55 AACGGCCACAAGTTCAGCGTGTCCGGCGAGGGCGAGGGCGATGCCACCTACGGCAAGCTG  
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60 ATCGAGCTGAAGGGCATCGACTTCAAGGAGGACGGCAACATCCTGGGGCACAAGCTGGAG

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5 TTCGTGACCGCCGCCGGGATCACTCTCGGCATGGACGAGCTGTACAAGTAAAGCGGCCCT  
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10 GAAACCAGCAGCGGCTATCCGCGCATCCATGCCCCGAAGTGCAGGAGTGGGGAGGCACG  
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15 ACTTACGGTAAATGGCCCGCTGGCTGACCGCCCAACGACCCCGCCCATTGACGTCAAT  
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20 GCGGTTTTGGCAGTACATCAATGGGCGTGGATAGCGGTTTGACTCACGGGATTTCGAAG  
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25 CATGGGGGACGGTGCCACCGGCCGCGCCATGGACGGGCGCGCCTGCTGCTGTTGCTGCT  
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30 GGAGGCCGACGACGCGGTGTGCCGTGCGCTACGGCTACTACCAGGATGAGACGACTGG  
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35 ACCCCCAGAGGGCTCGGACAGCACAGCCCCAGCACCCAGGAGCCTGAGGCACCTCCAGA  
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40 GGGGGGAATGAAAGACCCACCTGTAGGTTTGGCAAGCTAGCTTAAGTAACGCCATTTTG  
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45 GTTTCTAGAGAACCATCAGATGTTTCCAGGGTGCCCCAAGGACCTGAAATGACCCTGTGC  
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50 TGGGGGCTCGTCCGGGATCGGGAGACCCCTGCCAGGGACCACCGACCCACCGGGGAG  
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55 ATGCGGTGTGAAATACCGCACAGATGCGTAAGGAGAAAATACCGCATCAGGCGCTCTTCC  
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60 AACCCGACAGGACTATAAAGATACAGGCGTTTTCCCCCTGGAAGCTCCCTCGTGCCTCT

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 5 AGGATTAGCAGAGCGAGGTATGTAGGCGGTGCTACAGAGTCTTGAAGTGGTGGCCTAAC  
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 15 AGAAGTGGTCTGCAACTTTATCCGCTCCATCCAGTCTATTAATTGTTGCCGGGAAGCT  
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 20 TCTCTTACTGTCTATGCCATCCGTAAGATGCTTTTCTGTGACTGGTGAAGTACTCAACCAAG  
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 25 AGGCAAAATGCCGCAAAAAAGGGAATAAGGGCGACACGGAAATGTTGAATACTCATACTC  
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 30 TCCCCGGAGACGGTCACAGCTTGTCTGTAAGCGGATGCCGGGAGCAGACAAGCCCGTCAG  
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 35 TGACGTCAATAATGACGTATGTTCCCATAGTAACGCCAATAGGGACTTTCCATTGACGTC  
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 40 GATTTCCAAGTCTCCACCCCATTTGACGTCAATGGGAGTTTGTGTTGGCACCAAAATCAAC  
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 45 TTTCAATTTGGGGGCTCGTCCGGGATTTGGGAGACCCCTGCCAGGGACCACCGACCCACCA  
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 50 AGGATATGTGGTTCTGGTAGGAGACGAGAACCTAAAACAGTTCCCGCCTCCGTCTGAATT  
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 TACCACTCCCTTAAGTTTACCTTAGGTCACTGGAAAGATGTCGAGCGGATCGCTCACAA  
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## PLTRI xP (SEQ ID No 58)

GCTAGCATAACTTCGTATAATGTATGCTATACGAAGTTATTCTAGAGAACCATCAGATGT  
 60 TTCCAGGGTGCCCCAAGGACCTGAAATGACCCGTGTCCTTATTTGAACTAACCAATCAGT

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 5 GACCCCTGCCAGGGACCACCGACCCACCACCGGGAGGTAAGCTGGCTGCCTCGCGCGTT  
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 10 ATGCGTAAGGAGAAAATACCGCATCAGGCGCTCTTCCGCTTCCCTCGCTCACTGACTCGCT  
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 15 CCAGGCGTTTCCCCCTGGAAGCTCCCTCGTGCCTCTCTGTTCCGACCCTGCCGCTTAC  
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 20 AGGCGGTGCTACAGAGTTCTTGAAGTGGTGGCTAACTACGGCTACACTAGAAGGACAGT  
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 25 CTAGATCCTTTTAAATTAATAATGAAGTTTTAAATCAATCTAAAGTATATATGAGTAAAC  
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 TCGTTTCATCCATAGTTGCCTGACTCCCCGTGCTGTAGATAACTACGATACGGGAGGGCTT  
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 30 CGCTCCATCCAGTCTATTAATTGTTGCGGGAAGCTAGAGTAAGTAGTTCGCCAGTTAA  
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 AGTGTATCACTCATGGTTATGGCAGCACTGCATAATTCTTCTACTGTATGCCATCCGT  
 35 AAGATGCTTTTCTGTGACTGGTGAAGTCAACCAAGTCATTCTGAGAATAGTGTATGCG  
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 40 AATAAGGGCGACACGGAAATGTTGAATACTCATACTCTTCTTTTCAATATTATTGAAG  
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 45 GCGGGCTTCTGCCTCTTAGACCACTCTACCCTATTCCCCACACTACCGGAGCCAAAGCC  
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### LTR plasmid (SEQ ID No 59)

GCTAGCTTAAGTAACGCCATTTTGCAAGGCATGGAAAAATACATAACTGAGAATAGAGAA  
 50 GTTCAGATCAAGGTCAGGAACAGATGGAACAGCTGAATATGGGCCAAACAGGATATCTGT  
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 AAACAGGATATCTGTGGTAAGCAGTTCCTGCCCCGGCTCAGGGCCAAGAACAGATGGTCC  
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 55 CTGTTTCGCGCGCTTCTGCTCCCCGAGCTCAATAAAAGAGCCCACAACCCCTCACTCGGGG  
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5 AAATACCGCATCAGGCGCTCTCCGCTTCCTCGCTCACTGACTCGCTGCGCTCGGTCTGTT  
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